

Db	781	AGACACGGTGACCGAGATAAGCTCCTCTCTCCAGGCCACCCGGCCAACTCCTTTCTACT	840
Qy	978	ACCCGGGGGTGAAGGCCCTTCCTCCCATCGCCAGGGTGACACTGGTGCGGCTCGCACAGA	1037
Db	841	ACCCGGGGGTGAAGGCCCTTCCTCCCATCGCCAGGGTGACACTGGTGCGGCTCGCACAGA	900
Qy	1038	GCCCCAGGGCCCTTCATCCCTCCCGGCCCAAGTCTCTGCCAGAGGACAATGAGATGTGTAG	1097
Db	901	GCCCCAGGGCCCTTCATCCCTCCCGGCCCAAGTCTCTGCCAGAGGACAATGAGATGTGTAG	960
Qy	1098	ACAGCGCCTCAGTTCCAGAAACCGCGCTGGAGTCTCGAGGTTCTCCCTGTGTGCTGCTGGG	1157
Db	961	ACAGCGCCTCAGTTCCAGAAACCGCGCTGGAGTCTCGAGGTTCTCCCTGTGTGCTGCTGGG	1020
Qy	1158	GACTGTGCGGAGGCCACTGTGGGAGGCTCGGAGCCAAAGACGAGACTCGGTACGTCGCGG	1217
Db	1021	GACTGTGCGGAGGCCACTGTGGGAGGCTCGGAGCCAAAGACGAGACTCGGTACGTCGCGG	1080
Qy	1218	TCAGAGCCGCCAACACGAGGAGCCCTCGCCCGAGCTCGAAGAGAGAGGCTGAGTGGGTCC	1277
Db	1081	TCAGAGCCGCCAACACGAGGAGCCCTCGCCCGAGCTCGAAGAGAGAGGCTGAGTGGGTCC	1140
Qy	1278	CTGATAACTGCGTCTTAAGACACGAGCCCGAGCCCTTGGGGCCCCCGGAGGCATATGGGG	1337
Db	1141	CTGATAACTGCGTCTTAAGACACGAGCCCGAGCCCTTGGGGCCCCCGGAGGCATATGGGG	1200
Qy	1338	TGTCTGGGGGCTCTGTGTGCAGGCTCATGTGTGAGGCGGCGGAGGCGCATGTTCGGG	1397
Db	1201	TGTCTGGGGGCTCTGTGTGCAGGCTCATGTGTGAGGCGGCGGAGGCGCATGTTCGGG	1260
Qy	1398	CTGCTCTTGACCGCGGTGAGGCGCGCGACCATCTCTGCACTCAAGGGGCCCTCTGGTGG	1457
Db	1261	CTGCTCTTGACCGCGGTGAGGCGCGCGACCATCTCTGCACTCAAGGGGCCCTCTGGTGG	1320
Qy	1458	CCGGCACGGGCAATGGGAAACAGCCTCTCTCTTTCCCAACCTTGCTTCTTAGGGGGCCCC	1517
Db	1321	CCGGCACGGGCAATGGGAAACAGCCTCTCTCTTTCCCAACCTTGCTTCTTAGGGGGCCCC	1380
Qy	1518	GTCTCCCGTCTGCTCTACGCTTCCTCTCTGCAAGTAAAGTTCATCCCCAAGGCTCCAG	1577
Db	1381	GTCTCCCGTCTGCTCTACGCTTCCTCTCTGCAAGTAAAGTTCATCCCCAAGGCTCCAG	1440
Qy	1578	CTACTCTAAATATGTCTCTTATAAGTTATTGTCTGCCAGGAGATTGTCTTTCATCGT	1637
Db	1441	CTACTCTAAATATGTCTCTTATAAGTTATTGTCTGCCAGGAGATTGTCTTTCATCGT	1500
Qy	1638	CCAGGGGCTGGCTCCACGTGGTTGCAAGTACCTCAGACCTGGTGTCTAGGCTGTGCT	1697
Db	1501	CCAGGGGCTGGCTCCACGTGGTTGCAAGTACCTCAGACCTGGTGTCTAGGCTGTGCT	1560
Qy	1698	GAGCCACATCTCCGAGGGCGCATCCAAGCGGGGCCACTTGAGAAAGTGAATAAATGGGG	1757
Db	1561	GAGCCACATCTCCGAGGGCGCATCCAAGCGGGGCCACTTGAGAAAGTGAATAAATGGGG	1620
Qy	1758	CGGTTTTCGGAAGCGTCAGTGTGTTTCCATGTTATGGATCTCTCTCGGTTTGATAAAGACTA	1817
Db	1621	CGGTTTTCGGAAGCGTCAGTGTGTTTCCATGTTATGGATCTCTCTCGGTTTGATAAAGACTA	1680
Qy	1818	TCCTCTGTTGCTCAAAAAAATAAAAAA 1840	
Db	1681	TCCTCTGTTGCTCAAAAAAATAAAAAA 1703	

RESULT 13

AAH34981

ID AAH34981 standard; cDNA: 1848 BP.

XX
XX
XXXXXAC
AAH

AC YY

100C-333-20: 33

DT 03-SEP-2001 (first entry)

Human; colon cancer; colon cancer antigen; diagnosis; detection; colorectal carcinoma; ss.

Homo sapiens.
 WO200122920-A2.
 05-APR-2001.
 28-SEP-2000; 2000WO-US26524.
 29-SEP-1999; 99US-0157137.
 03-NOV-1999; 99US-0163280.
 (HUMA-) HUMAN GENOME SCI INC.
 Ruben SM, Barash SC, Birse CE, Rosen CA;
 WPI: 2001-235357/24;
 P-PSDB: AAG75576.
 Nucleic acids encoding 4277 human colon cancer-associated polypeptides,
 useful for preventing, diagnosing and/or treating colorectal cancers -
 Claim 1; Page 3549; 9803pp; English.

Sequence 1848 BP; 324 A; 601 C; 578 G; 340 T; 5 other; XX

```
Query Match      91.3%; Score 1680.8; DB 22; Length 1848;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 1679; Conservative 3; Mismatches 1; Indels 0;
```

158	QY	AGCCCTGCAGCATCGAAGACAGAGGAACTTGAGGCTCTATTGGCCGGCCCGGGCGCCG	217
159	QY		
160	QY		
161	QY		
162	QY		
163	QY		
164	QY		
165	QY		
166	QY		
167	QY		
168	QY		
169	QY		
170	QY		
171	QY		
172	QY		
173	QY		
174	QY		
175	QY		
176	QY		
177	QY		
178	QY		
179	QY		
180	QY		
181	QY		
182	QY		
183	QY		
184	QY		
185	QY		
186	QY		
187	QY		
188	QY		
189	QY		
190	QY		
191	QY		
192	QY		
193	QY		
194	QY		
195	QY		
196	QY		
197	QY		
198	QY		
199	QY		
200	QY		
201	QY		
202	QY		
203	QY		
204	QY		
205	QY		
206	QY		
207	QY		
208	QY		
209	QY		
210	QY		
211	QY		
212	QY		
213	QY		
214	QY		
215	QY		
216	QY		
217	QY		
218	QY		
219	QY		
220	QY		
221	QY		
222	QY		
223	QY		
224	QY		
225	QY		
226	QY		
227	QY		
228	QY		
229	QY		
230	QY		
231	QY		
232	QY		
233	QY		
234	QY		
235	QY		
236	QY		
237	QY		
238	QY		
239	QY		
240	QY		
241	QY		
242	QY		
243	QY		
244	QY		
245	QY		
246	QY		
247	QY		
248	QY		
249	QY		
250	QY		
251	QY		
252	QY		
253	QY		
254	QY		
255	QY		
256	QY		
257	QY		
258	QY		
259	QY		
260	QY		
261	QY		
262	QY		
263	QY		
264	QY		
265	QY		
266	QY		
267	QY		
268	QY		
269	QY		
270	QY		
271	QY		
272	QY		
273	QY		
274	QY		
275	QY		
276	QY		
277	QY		
278	QY		
279	QY		
280	QY		
281	QY		
282	QY		
283	QY		
284	QY		

QY	458	ACGGCTTCCCAAGCAGTACCCCTGTTCCGCCCCTCTGCGCAGTGGTCTTCGCTGCTG 517	Db	1513	CTCCTCCTCTGAGGATAAGTCAATGAGGCTCCAGCTACTCTAAATTTATGTTCTCC 1572
Db	433	ACGGCTTCCCAAGCAGTACCCCTGTTCCGCCCCTCTGCGCAGTGGTCTTCGCTGCTG 492	QY	1598	TTATAAGTTATTGCTCTCCAGGAGATTGCTTTCATGTCAGGGGCTTGGCTCCACG 1657
QY	518	GGGCGCGCATAGCTCCGACTACAGCATGTGAGGAAGAACACAGTACGTACGTACGGG 577	Db	1573	TTATAAGTTATTGCTCTCCAGGAGATTGCTTTCATGTCAGGGGCTTGGCTCCACG 1632
Db	493	GGGCGCGCATAGCTCCGACTACAGCATGTGAGGAAGAACACAGTACGTACGTACGGG 552	QY	1658	TGTTTCAGATACCTCAGACCTGGTCTCTAGGCTGTGCTGAGCCACCTCTCCCGAGGGC 1717
QY	578	CTGCGGACTTTCGAGAGCGCGGAGGCTGGCGCTGATGAAGAGATCGAGGCGGG 637	Db	1633	TGTTTCAGATACCTCAGACCTGGTCTCTAGGCTGTGCTGAGCCACCTCTCCCGAGGGC 1692
Db	553	CTGCGGACTTTCGAGAGCGCGGAGGCTGGCGCTGATGAAGAGATCGAGGCGGG 612	QY	1718	GCATCAAGCGGGGCGACCTGAGAAAGTGAATAAATGGGCGGTTTCGGAAGCGTCAGTG 1777
QY	638	GGGAGGCGCTGCAGAGCGTGCAGAGGTGTTTTCGCGCGCCCGCTCCCGACGCGCAC 697	Db	1693	GCATCAAGCGGGGCGACCTGAGAAAGTGAATAAATGGGCGGTTTCGGAAGCGTCAGTG 1752
Db	613	GGGAGGCGCTGCAGAGCGTGCAGAGGTGTTTTCGCGCGCCCGCTCCCGACGCGCAC 672	QY	1778	TTTCCATGTTATGATCTCTCTCGGTTTGAATAAAGACTACTCTCTGCTCAAAAAA 1837
QY	698	GGGAGAGCTGCGGAGCTGGAGGTGACAGCGAGGACTCGCTGTGCTCTGTTTGGTG 757	Db	1753	TTTCCATGTTATGATCTCTCTCGGTTTGAATAAAGACTACTCTCTGCTCAAAAAA 1812
Db	673	GGGAGAGCTGCGGAGCTGGAGGTGACAGCGAGGACTCGCTGTGCTCTGTTTGGTG 732	QY	1838	AAA 1840
QY	758	CGATCTGCGGAGCGGCGGCTGGTCTGTTGGGCTGACAGCTTGGACCTGTGCGAGCGG 817	Db	1813	AAA 1815
Db	733	CGATCTGCGGAGCGGCGGCTGGTCTGTTGGGCTGACAGCTTGGACCTGTGCGAGCGG 792	RESULT 14		
QY	818	GACGTTGGCGGAACAGGCGGCTGGACCTGTACCCCTACGACGCGGCGGACGACG 877	AAK94182		
Db	793	GACGTTGGCGGAACAGGCGGCTGGACCTGTACCCCTACGACGCGGCGGACGACG 852	ID AAK94182 standard; cDNA; 1669 BP.		
QY	878	GGCTTACCTTCTCTCTCCCAACTTCGCCACCACTCCCGCAGGACGCGTACCGAGATA 937	XX AAK94182;		
Db	853	GGCTTACCTTCTCTCTCCCAACTTCGCCACCACTCCCGCAGGACGCGTACCGAGATA 912	AC		
QY	938	ACGCTCTCTCTCTCCAGCACCGCGCAACTCTCTACTACCGCGGCTGAAGCGCTG 997	XX		
Db	913	ACGCTCTCTCTCTCCAGCACCGCGCAACTCTCTACTACCGCGGCTGAAGCGCTG 972	DT		
QY	998	CCTCCCATCGGAGTGCACACTGTGTGGCTGGACAGAGCCCGGCTTCATCCCT 1057	DE		
Db	973	CCTCCCATCGGAGTGCACACTGTGTGGCTGGACAGAGCCCGGCTTCATCCCT 1032	XX		
QY	1058	CCCGCCCGAGTCTGCCAGCAGGACAAATGAGATTGATAGAGCGCTCAGTTCAGAA 1117	OS		
Db	1033	CCCGCCCGAGTCTGCCAGCAGGACAAATGAGATTGATAGAGCGCTCAGTTCAGAA 1092	XX		
QY	1118	ACGCGCTGGAGTCTCCCTGTGCTGCTGCGGAGTCTGCGGAGGCTGCGGAGGCTGCT 1177	PN		
Db	1093	ACGCGCTGGAGTCTCCCTGTGCTGCTGCGGAGTCTGCGGAGGCTGCTGCGGAGGCT 1152	XX		
QY	1178	GGGAGGCTCGGACCAAGAGCAGGACTCGCTACGTCCGGGTCCAGCGCCGCCAACACGG 1237	PD		
Db	1153	GGGAGGCTCGGACCAAGAGCAGGACTCGCTACGTCCGGGTCCAGCGCCGCCAACACGG 1212	PF		
QY	1238	AGCCCTGCGCGAGTCTGAGGAGGCTGAGTGGCTCCCTGATTAACGTGCTCTAAGAC 1297	XX		
Db	1213	AGCCCTGCGCGAGTCTGAGGAGGCTGAGTGGCTCCCTGATTAACGTGCTCTAAGAC 1272	PR		
QY	1298	CAGAGCCCGCAGCCCTCGGCGCCCGGAGGCTGCTGCGGCTCTCTGTCAG 1357	PR		
Db	1273	CAGAGCCCGCAGCCCTCGGCGCCCGGAGGCTGCTGCGGCTCTCTGTCAG 1332	PR		
QY	1358	GCTCATGCTGCGAGGCGCGGAGGACAGGGGTTTCGCGCTGCTCTGACCGCGGTGAG 1417	XX		
Db	1333	GCTCATGCTGCGAGGCGCGGAGGACAGGGGTTTCGCGCTGCTCTGACCGCGGTGAG 1392	PI		
QY	1418	GCGCGCGCAGCATTCTGCTGATGAAGGCGCTCTGGTGGCGGCGACGGGCTTGGGAA 1477	XX		
Db	1393	GCGCGCGCAGCATTCTGCTGATGAAGGCGCTCTGGTGGCGGCGACGGGCTTGGGAA 1452	DR		
QY	1478	CAGCTCTCTCTTCCCACTTCTTCTAGGGGCGCGGCTCTGCTGCTCTCAGC 1537	XX		
Db	1453	CAGCTCTCTCTTCCCACTTCTTCTAGGGGCGCGGCTCTGCTGCTCTCAGC 1512	PT		
QY	1538	CTCTCTCTCTGCGAGGATAAGTCAATGAGGCTCCAGCTACTCTAAATTTATGTTCTCC 1597	PS		

Human full-length cDNA, SEQ ID NO: 2726.

Human; full length cDNA; cDNA synthesis; oligo-capping; ss.

Homo sapiens.

EP1130094-A2.

05-SEP-2001.

07-JUL-2000; 2000EP-0114089.

08-JUL-1999; 99JP-0194486.

11-JAN-2000; 2000JP-0118774.

02-MAY-2000; 2000JP-0183765.

(HELI-) HELIX RES INST.

Ota T, Nishikawa T, Isogai T, Hayashi K, Ishii S, Kawai Y, Wakamatsu A, Sugiyama T, Nagai K, Kojima S, Otsuki T, Koga H;

WPI: 2001-524255/58.

P-PSDB; AAM93266.

830 Primers useful for synthesizing full length cDNA clones and their use in genetic manipulation -

Claim 8; SEQ ID NO 2726; 1380pp + sequence listing; English.

The invention relates to primers for synthesizing full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5' and 3'-ends of the cDNA molecules have been determined. Primers for synthesizing the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesized by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a full length human cDNA of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in CD-ROM format directly from EPO.

QY 1090 GATTGTAGACAGCGCTCAGTTCAGAAACGCGCGCTGACTGCGAGGTCTCCCTGTGGTC 1149
|||||
Db 236 GATTGTAGACAGCGCTCAGTTCAGAAACGCGCGCTGACTGCGAGGTCTCCCTGTGGTC 295
|||||
QY 1150 GTCTGGGACTGTGGGAGGCGCTGTGGAGGCTCGGAGCAACAGAGGACTCGCTA 1209
|||||
Db 296 GTCTGGGACTGTGGGAGGCGCTGTGGAGGCTCGGAGCAACAGAGGACTCGCTA 355
|||||
QY 1210 CGTCCGGTCCAGCGCGCAACAGCGGAGCGCTGCGCGGAGCT 1254
|||||
Db 356 CGCCCGGGTCCAGCGCGCAACAGCGGAGCGCTGCGCGGAGCT 400
|||||

RESULT 4

US-09-022-238-1
; Sequence 1, Application US/0902238
; Patent No. 6177244
; GENERAL INFORMATION:
; APPLICANT: Sytkowski, Arthur J. and Yang, Meiheng
; TITLE OF INVENTION: A novel NPG-1 Gene that is differentially expressed in prosta
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LAHIVE & COCKFIELD, LLP
; STREET: 28 State Street
; CITY: Boston
; STATE: Massachusetts
; COUNTRY: USA
; ZIP: 02109

COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/022,238
; FILING DATE:
; CLASSIFICATION:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: US 08/644,326
; FILING DATE: 10-MAY-1996

ATTORNEY/AGENT INFORMATION:
; NAME: Maravic-Magovcevic, Ivana
; REGISTRATION NUMBER: P-43,338
; REFERENCE/DOCKET NUMBER: NER-262CP
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617)227-7400
; TELEFAX: (617)742-4214
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 400 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 3..398
US-09-022-238-1

Query Match 18.48; Score 339; DB 4; Length 400;
Best Local Similarity 96.3%; Pred. No. 5.3e-62;
Matches 390; Conservative 0; Mismatches 10; Indels 5; Gaps 4;

QY 850 GTACCCCTACACCGCGGAGCGGAGTTCACCTTCTCCCTCCCGCAACTTCGGCAC 909
|||||
Db 1 GTACCCCTACACCGCGGAGGAC-GACAGCGGCTTACCTTCTCCCTCCCGCAACTTCGGCAC 59
|||||
QY 910 CATCCCGAGGACAGGTGACCGAGATAAAGTCTCTCTCCAGCCAGCGGCGCAACTC 969
|||||
Db 60 CATCCCGCA-GACACGCTGACCGAGATACGTCTCTCTCCAGCCAGCGGCGCAACTC 118
|||||
970 CTTCCTACCCCGGCTGAAGGCCCTGCCCTCCCATCGCCAGGCTGACACTGTGCGGCT 1029

Db 119 CTCTACTACCGCGGCTGAGGCC--TGCTCCATCGCCAGGGTGACACTGGTGGGCT 176
|||||
QY 1030 GCACAGAGCCCGAGGCGCTTATCCCTCCCGCCAGTCTGCCAGGAGGACAATGA 1089
|||||
Db 177 GCACAGAGCCCGAGGCGCTTATCCCTCCCGCCAGTCTGCCAGGAGGACAATGA 235
|||||
QY 1090 GATTGTAGACAGCGCTCAGTTCAGAAACGCGCGCTGACTGCGAGGTCTCCCTGTGGTC 1149
|||||
Db 236 GATTGTAGACAGCGCTCAGTTCAGAAACGCGCGCTGACTGCGAGGTCTCCCTGTGGTC 295
|||||
QY 1150 GTCTGGGACTGTGGGAGGCGCTGTGGAGGCTCGGAGCAACAGAGGACTCGCTA 1209
|||||
Db 296 GTCTGGGACTGTGGGAGGCGCTGTGGAGGCTCGGAGCAACAGAGGACTCGCTA 355
|||||
QY 1210 CGTCCGGTCCAGCGCGCAACAGCGGAGCGCTGCGCGGAGCT 1254
|||||
Db 356 CGCCCGGGTCCAGCGCGCAACAGCGGAGCGCTGCGCGGAGCT 400
|||||

RESULT 5

US-08-799-173A-15
; Sequence 15, Application US/08799173A
; Patent No. 5871969
; GENERAL INFORMATION:
; APPLICANT: HASTINGS, GREGG,
; APPLICANT: PATRICK J. DILLON

; TITLE OF INVENTION: HUMAN NEURONAL ATTACHMENT FACTOR-1
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HUMAN GENOME SCIENCES, INC.
; STREET: 9410 KEY WEST AVENUE
; CITY: ROCKVILLE
; STATE: MD
; COUNTRY: USA
; ZIP: 20850

COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/799,173A
; FILING DATE: 11-FEB-1997
; CLASSIFICATION: 536
; ATTORNEY/AGENT INFORMATION:
; NAME: BROOKES, ANDERS A.
; REGISTRATION NUMBER: 36,373
; REFERENCE/DOCKET NUMBER: PF226
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (301) 309-8504
; TELEFAX: (301) 309-8512
; INFORMATION FOR SEQ ID NO: 15:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 506 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-799-173A-15

Query Match 16.68; Score 304.8; DB 2; Length 506;
Best Local Similarity 85.5%; Pred. No. 6.9e-55;
Matches 371; Conservative 0; Mismatches 59; Indels 4; Gaps 4;

QY 293 CGGTGTGTAAGAAACCCAGCGCGCGCGCGCTGGCAAGGCCCTCTGGCTCTCTC 352
|||||
Db 6 CGCANAGNNAAACCCAGCGCGCGCGCTGGCAAGGCCCTCTGGCTCTCTC 65
|||||
QY 353 CTGCCACTCTCGCGCGCGCGCGCGCTCTTTGGGGAGAGTCCATCTGTTCGCCGAGA 412
|||||
Db 66 CTGCCACTCTCGCGCGCGCGC-ACCAGCCTCTTTGGGGAGAGTCCATCTTTTCGCCGAGA 124
|||||

Result No.	Query			ID	Description
	Score	Match	Length		
1	1692.4	92.0	1779	4	US-09-371-696-1
2	1100.2	59.8	1105	2	US-08-799-173A-1
3	343.8	18.7	400	1	US-08-644-326-1
4	339	18.4	400	4	US-09-022-238-1
5	304.8	16.6	506	2	US-08-799-173A-15
6	204.2	11.1	316	2	US-08-799-173A-16
7	204.2	11.1	316	2	US-08-799-173A-17
8	71.6	3.9	4029	1	US-07-862-021B-9
9	71.6	3.9	4029	1	US-08-313-288B-9
10	71.6	3.9	4029	5	PCT-US93-03164-9
11	62.2	3.4	3226	1	US-07-862-021B-11
12	62.2	3.4	3226	1	US-08-313-288B-11
13	62.2	3.4	3226	5	PCT-US93-03164-11
14	57.6	3.1	1155	4	US-08-818-112-12
15	57.6	3.1	1155	4	US-08-818-111-12
16	57.6	3.1	1155	4	US-09-056-556-12
17	57.2	3.1	1816	1	US-07-862-021B-13
18	57.2	3.1	1816	1	US-08-313-288B-13
19	57.2	3.1	1816	5	PCT-US93-03164-13
20	55.4	3.0	1105	2	US-08-799-173A-1
21	55	3.0	30001	1	US-08-125-468-1
22	55	3.0	30001	2	US-08-474-933-1
23	54.4	3.0	933	4	US-09-105-390-43
24	54.4	3.0	1008	4	US-09-105-390-59
25	54.4	3.0	2810	4	US-09-105-390-6
26	52.4	2.8	1779	4	US-09-371-696-1
27	51.8	2.8	8625	4	US-08-960-832-1

